

10/578797

SEQUENCE LISTING  
AP 26 Rec'd PCT/PTO 04 MAY 2006

<110> Foss, Francine M.  
Xie, Yong

<120> Treatment with Immunoregulatory T Cells

<130> 00398-155US1

<140> To Be Assigned

<141> 2006-05-03

<150> PCT/US2004/037201

<151> 2004-11-05

<150> US 60/517,624

<151> 2003-11-05

<160> 2

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 1574

<212> DNA

<213> Homo sapiens

<400> 1

```

gagtagagcc gatctcccg cccccgaggt tgctcctctc cgaggtctcc cgcgcccaa 60
gttctcccg ccccgaggtc tcccgcccc gaggtctccg cgccccgagg tctccgccc 120
caccatgcgg ctgggcagtc ctggactgct ctctctgctc ttcagcagcc ttcgagctga 180
tactcaggag aaggaagtca gagcgatggt aggcagcgac gtggagctca gctgcgcttg 240
ccctgaagga agccgttttg atttaaata tgtttacgta tattggcaaa ccagtgaagtc 300
gaaaaccgtg gtgacctacc acatcccaca gaacagctcc ttggaaaacg tggacagccg 360
ctaccggaac cgagccctga tgtcaccggc cggcatgctg cggggcgact tctccctgcg 420
cttgttcaac gtcaccccc aggacgagca gaagtttcac tgcctggtgt tgagccaatc 480
cctgggattc caggaggttt tgagcggtga ggttacactg catgtggcag caaacttcag 540
cgtgcccgtc gtcagcgccc cccacagccc ctcccaggat gagctcacct tcacgtgtac 600
atccataaac ggctacccca ggcccaacgt gtactggatc aataagacgg acaacagcct 660
gctggaccag gctctgcaga atgacaccgt cttcttgaac atgcggggct tgtatgacgt 720
ggtcagcggt ctgaggatcg cacggacccc cagcgtgaac attggctgct gcatagagaa 780
cgtgcttctg cagcagaacc tgactgtcgg cagccagaca ggaaatgaca tcggagagag 840
agacaagatc acagagaatc cagtcagtac cggcgagaaa aacgcggcca cgtggagcat 900
cctggctgtc ctgtgcctgc ttgtggtcgt ggcggtggcc ataggctggg tgtgcaggga 960
ccgatgcctc caacacagct atgcaggtgc ctgggctgtg agtccggaga cagagctcac 1020
tggccacggt tgaccggagc tcaccgccc gagcgtggac agggcttccg tgagacgcca 1080
ccgtgagagg ccagggtggca gcttgagcat ggactcccag actgcagggg agcacttggg 1140
gcagccccc gaaggaccac tgcctggatc cagggagaa ctgctggcgt tggctgtgat 1200
cctggaatga ggccctttca aaagcgtcat ccacaccaa ggcaaattgc cccaagtga 1260
tgggctcccc gctgtcactg ccagtcaccc acaggaaggg actggtgatg ggctgtctct 1320
accggagcgt tgcgggattc agcaccaggc tcttcccagt accccagacc cactgtgggt 1380
cttcccgtgg gatgcgggat cctgagaccg aaggtgtttt ggtttaaaaa gaagactggg 1440
cgtccgctct tccaggacgg cctctgtgct gctggggtca cgcgaggctg tttgcagggg 1500
acacggtcac aggagctctt ctgcccgtga cgctcccaac ctgcctccc cccggaagcc 1560
acaggacca ctca

```

<210> 2

<211> 302

<212> PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 2

```

Met Arg Leu Gly Ser Pro Gly Leu Leu Phe Leu Leu Phe Ser Ser Leu
 1           5           10           15
Arg Ala Asp Thr Gln Glu Lys Glu Val Arg Ala Met Val Gly Ser Asp
      20           25           30
Val Glu Leu Ser Cys Ala Cys Pro Glu Gly Ser Arg Phe Asp Leu Asn
      35           40           45
Asp Val Tyr Val Tyr Trp Gln Thr Ser Glu Ser Lys Thr Val Val Thr
      50           55           60
Tyr His Ile Pro Gln Asn Ser Ser Leu Glu Asn Val Asp Ser Arg Tyr
65           70           75           80
Arg Asn Arg Ala Leu Met Ser Pro Ala Gly Met Leu Arg Gly Asp Phe
      85           90           95
Ser Leu Arg Leu Phe Asn Val Thr Pro Gln Asp Glu Gln Lys Phe His
      100          105          110
Cys Leu Val Leu Ser Gln Ser Leu Gly Phe Gln Glu Val Leu Ser Val
      115          120          125
Glu Val Thr Leu His Val Ala Ala Asn Phe Ser Val Pro Val Val Ser
      130          135          140
Ala Pro His Ser Pro Ser Gln Asp Glu Leu Thr Phe Thr Cys Thr Ser
145           150           155           160
Ile Asn Gly Tyr Pro Arg Pro Asn Val Tyr Trp Ile Asn Lys Thr Asp
      165          170          175
Asn Ser Leu Leu Asp Gln Ala Leu Gln Asn Asp Thr Val Phe Leu Asn
      180          185          190
Met Arg Gly Leu Tyr Asp Val Val Ser Val Leu Arg Ile Ala Arg Thr
      195          200          205
Pro Ser Val Asn Ile Gly Cys Cys Ile Glu Asn Val Leu Leu Gln Gln
      210          215          220
Asn Leu Thr Val Gly Ser Gln Thr Gly Asn Asp Ile Gly Glu Arg Asp
225           230           235           240
Lys Ile Thr Glu Asn Pro Val Ser Thr Gly Glu Lys Asn Ala Ala Thr
      245          250          255
Trp Ser Ile Leu Ala Val Leu Cys Leu Leu Val Val Val Ala Val Ala
      260          265          270
Ile Gly Trp Val Cys Arg Asp Arg Cys Leu Gln His Ser Tyr Ala Gly
      275          280          285
Ala Trp Ala Val Ser Pro Glu Thr Glu Leu Thr Gly His Val
      290          295          300

```